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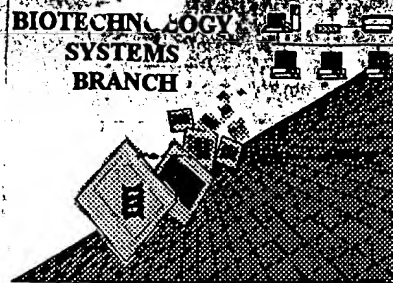
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RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/697,206

Source: C. PE

Date Processed by STIC: 11/8/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/697,206

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid-number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 8 Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of <213>Organism (NEW RULES) Sequence(s) are missing this mandatory field or its response.
- 12 Use of <220>Feature (NEW RULES) Sequence(s) are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000
 TIME: 10:45:55

Input Set : A:\Seqlist.txt
 Output Set: N:\CRF3\11082000\1697206.raw

Does Not Comply
 Corrected Diskette Needed

4 <110> APPLICANT: Daniel E.H. Afar
 5 Arthur B. Raitano
 6 Rene S. Hubert
 7 Steve Chappell Mitchell
 8 Aya Jakobovits
 10 <120> TITLE OF INVENTION: NOVEL GENE UPREGULATED IN CANCERS OF THE
 11 PROSTATE
 13 <130> FILE REFERENCE: 129.21-US-U1
 15 <140> CURRENT APPLICATION NUMBER: US/09/697,206
 15 <141> CURRENT FILING DATE: 2000-10-26
 15 <150> PRIOR APPLICATION NUMBER: 60/162,364
 16 <151> PRIOR FILING DATE: 1999-10-28
 18 <160> NUMBER OF SEQ ID NOS: 26
 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 3585
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Homo Sapiens
 27 <400> SEQUENCE: 1
 28 ctttttggga tcactgctgg ggccaccggg gccaaagctag gctcggatga gaaggagttg 60
 29 atcctgctgt tctggaaagt cgtggatctg gccaaacaaga aggtgggaca gttgcacgaa 120
 30 gtgctagtta gaccggatca gttggaaactg acggaggact gcaagaaga aactaaaata 180
 31 gacgtcgaaa gctgtctctc ggcgtcgcag ctggaccaag cctcccgaca gtttaaccag 240
 32 tcagttagca atgaactgaa tattggagta gggacttctt tctgtctctg tactgatggg 300
 33 cagcttcatg tcaggcaaat cctgcattct gaggcttcca agaagaatgt actattacct 360
 34 gaatgcttct attccttttt tgatcttcga aaagaattca agaatgttg ccttggttca 420
 35 cctgatattg acaaaactga cgttgccaca atgacagagt atttaaattt tgagaagagt 480
 36 agttcagttc ctcgatatgg agcctctcaa gttgaagata tggggaatat aatttttagca 540
 37 atgatttcag agccttataa tcacaggttt tcagatccag agagagtga ttacaagttt 600
 38 gaaagtggaa cttgcagcaa gatggaactt attgatgata acaccgtagt cagggcacga 660
 39 ggtttaccat ggcagttctc agatcaagat attgcaagat tcttcaaagg actcaatatt 720
 40 gccaaaggag gtgcagcact ttgtctgaat gctcagggtc gaaggaaagg agaagctctg 780
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 43 ggtacttcca atgaggttag ccagtttctc tccaaggaaa atcaagtcatt tgttcgcatg 960
 44 cgggggctcc ctttcacggc cacagctgaa gaagtgggtg ctttctcttg acagcaltgc 1020
 45 cctattactg ggggaaagga aggcattctc ttgtctcact acccagatgg taggccaaca 1080
 46 ggggaacgct ttgtctctct tgctgtgag gaatatgcac agaattgcgtt gagggaagcat 1140
 47 aaagacttgt tgggtaaaag atacattgaa ctcttcaggga gcacagcagc tgaagtccag 1200
 48 caggtgtgta atcgattctc ctccggccct ctcatctcac ttccaaaccc tccattattt 1260
 49 ccagtactac ctacgcaatt tgtgcccctt acaaatgtta gagactgtat acgctctoga 1320
 50 ggtcttccct atgcagccac aattgaggac atcctggatt tcttggggga gttcgccaca 1380
 51 gatattcgta ctcatgggtt tcacatgggt ttgaatcacc agggccgccc atcaggagat 1440
 52 gcctttatcc agatgaagtc tgcggacaga gcatttatgg ctgcacagaa gtgtcataaa 1500
 53 aaaaacatga aggacagala tgttgaagtc ttccagtgtt cagctgagga gatgaacttt 1560
 54 gtgttaatgg ggggcaactt aaatcgaaat ggttatecco caccgcaatg cctgtctcct 1620
 55 cctcctaca catttccagc tctgtgtgca gttattccta cagaagctgc catttaccag 1680

pp 4-5

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000
 TIME: 10:45:55

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 Output Set: N:\CRF3\11082000\I697206.raw

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56 cccctctgtga ttttgaatcc acgagcactg cagccctcca cagcgtacta cccagcaggc 1740
57 actcagctct tcatgaacta cacagcgtac tatcccagcc cccaggttc gccaatagt 1800
58 cttggctact tccctacagc tgctaattctt agcgggtgtcc cccacacagc tggcacggty 1860
59 gtcagaatgc agggcctggc ctacaatact ggagttaagg aaattcttaa cttcttccaa 1920
60 ggttaccagt atgcaaccga ggaatggactt atacacacaa atgaccagge caggactcta 1980
61 cccaaagaat gggtttgtat ttaagggccc cagcagttag aacatcctca gaaaagaagt 2040
62 gtttgaaga tqtatggta tcttgaaacc tccaqacaca agaaaacttc tagcaaatc 2100
63 aggggaagtt tgcctacact cagcgtgcag tattttcagc aaacttgatt ggacaaacgg 2160
64 gcctgtgcct tatcttttgg tggagtgaag aaatttgagc tagtgaagcc aaatcgtaac 2220
65 ttacagcaag cagcagtcag catacctggc tctttgctga ttgcaaatag gcatttaaaa 2280
66 tgtgaatttg gaatcagatg tctccattac ttccagttaa agtggcatca taggtgtttc 2340
67 ctaagtttta agtcctggat aaaaactcca ccagtgctca ccatctccac catgaactct 2400
68 gtttaaggag cttcattttt gtatatcccc gctcttttct cttcatttcc ctgtctctg 2460
69 cataatcatg ccttctgtct agtaattcca agcataagat cttggaataa taaaatcacu 2520
70 atcttaggag aaagaataaa attgttattt tcccagtcct ttggccatga tgatatctta 2580
71 tgattaaaaa caaattaaat tttaaaacac ctgaagataa attagaagaa attgtgcacc 2640
72 cttccaaaaa catacaaagt ttaaaagttt ggatcttttt ctcagcaggt atcagttgta 2700
73 aataatgaat taggggccc aaatgcaaac gaaaaatgaa gcagctacat gtagttagta 2760
74 atttctagtt tgaactgtaa ttgaatatgt tggcttcata tgtattattt tatattgtac 2820
75 ttttttctatt attgatggtt tggactttta taagagaaat tccatagttt ttaatatccc 2880
76 agaagtgaga caatttgaac agtgtattct agaaaaaat acactaactg aacagaagt 2940
77 aatgcttata tatattatga tagcctttaa cctttttcct ctaattgcctt aactgtcaaa 3000
78 taattataac cttttaaagc ataggactat agtcagcatg ctgactgag aggtaaacac 3060
79 tgatgcaatt agaacaggta ctgatgtgt cagtgtttta cactatgttt agctgtgttt 3120
80 atgctataaa agtgcataat tagacactag ctagtactgc tgcctcatgt aactccaaag 3180
81 aaaacaggat ttcattaagt gcattgaatg tggatatctc tctaagttac tcatattgtc 3240
82 ctttgcttga atgcaatgcc gtgcagattt atgaggctgc tatttttatt tctgtgcat 3300
83 tactttaaca ctttaaaggg agaagcaaac atttcttctc tcagctgact ggcattggcc 3360
84 ctttaactgc aatagggaag aaaaaaaaaa ggtttggtg aaaaattggtg ataactggca 3420
85 cttaaagatg aaaagaaatt tctgtatact tgatgcctta agatgcccaa agctgcccaa 3480
86 agctctgaaa gactttaaga taggcagtaa tgcttactac aatactactg agtttttgta 3540
87 gagttaacat ttgataataa aacttgctg tttaatctca aaaaa 3585

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89 <210> SEQ ID NO: 2

90 <211> LENGTH: 517

91 <212> TYPE: PRT

92 <213> ORGANISM: Homo Sapiens

94 <400> SEQUENCE: 2

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95 Met Thr Glu Tyr Leu Asn Phe Glu Lys Ser Ser Ser Val Ser Arg Tyr
96 1 5 10 15
97 Gly Ala Ser Gln Val Glu Asp Met Gly Asn Ile Ile Leu Ala Met Ile
98 20 25 30
99 Ser Glu Pro Tyr Asn His Arg Phe Ser Asp Pro Glu Arg Val Asn Tyr
100 35 40 45
101 Lys Phe Glu Ser Gly Thr Cys Ser Lys Met Glu Leu Ile Asp Asp Asn
102 50 55 60
103 Thr Val Val Arg Ala Arg Gly Leu Pro Trp Gln Ser Ser Asp Gln Asp
104 65 70 75 80
105 Ile Ala Arg Phe Phe Lys Gly Leu Asn Ile Ala Lys Gly Gly Ala Ala
106 85 90 95

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000

TIME: 10:45:55

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\11082000\I697206.raw

```

107 Leu Cys Leu Asn Ala Gln Gly Arg Arg Asn Gly Glu Ala Leu Val Arg
108           100           105           110
109 Phe Val Ser Glu Glu His Arg Asp Leu Ala Leu Gln Arg His Lys His
110           115           120           125
111 His Met Gly Thr Arg Tyr Ile Glu Val Tyr Lys Ala Thr Gly Glu Asp
112           130           135           140
113 Phe Leu Lys Ile Ala Gly Gly Thr Ser Asn Glu Val Ala Gln Phe Leu
114 145           150           155           160
115 Ser Lys Glu Asn Gln Val Ile Val Arg Met Arg Gly Leu Pro Phe Thr
116           165           170           175
117 Ala Thr Ala Glu Glu Val Val Ala Phe Phe Gly Gln His Cys Pro Ile
118           180           185           190
119 Thr Gly Gly Lys Glu Gly Ile Leu Phe Val Thr Tyr Pro Asp Gly Arg
120           195           200           205
121 Pro Thr Gly Asp Ala Phe Val Leu Phe Ala Cys Glu Glu Tyr Ala Gln
122           210           215           220
123 Asn Ala Leu Arg Lys His Lys Asp Leu Leu Gly Lys Arg Tyr Ile Glu
124 225           230           235           240
125 Leu Phe Arg Ser Thr Ala Ala Glu Val Gln Gln Val Leu Asn Arg Phe
126           245           250           255
127 Ser Ser Ala Pro Leu Ile Pro Leu Pro Thr Pro Pro Ile Ile Pro Val
128           260           265           270
129 Leu Pro Gln Gln Phe Val Pro Pro Thr Asn Val Arg Asp Cys Ile Arg
130           275           280           285
131 Leu Arg Gly Leu Pro Tyr Ala Ala Thr Ile Glu Asp Ile Leu Asp Phe
132           290           295           300
133 Leu Gly Glu Phe Ala Thr Asp Ile Arg Thr His Gly Val His Met Val
134 305           310           315           320
135 Leu Asn His Gln Gly Arg Pro Ser Gly Asp Ala Phe Ile Gln Met Lys
136           325           330           335
137 Ser Ala Asp Arg Ala Phe Met Ala Ala Gln Lys Cys His Lys Lys Asn
138           340           345           350
139 Met Lys Asp Arg Tyr Val Glu Val Phe Gln Cys Ser Ala Glu Glu Met
140           355           360           365
141 Asn Phe Val Leu Met Gly Gly Thr Leu Asn Arg Asn Gly Leu Ser Pro
142           370           375           380
143 Pro Pro Cys Leu Ser Pro Pro Ser Tyr Thr Phe Pro Ala Pro Ala Ala
144 385           390           395           400
145 Val Ile Pro Thr Glu Ala Ala Ile Tyr Gln Pro Ser Val Ile Leu Asn
146           405           410           415
147 Pro Arg Ala Leu Gln Pro Ser Thr Ala Tyr Tyr Pro Ala Gly Thr Gln
148           420           425           430
149 Leu Phe Met Asn Tyr Thr Ala Tyr Tyr Pro Ser Pro Pro Gly Ser Pro
150           435           440           445
151 Asn Ser Leu Gly Tyr Phe Pro Thr Ala Ala Asn Leu Ser Gly Val Pro
152           450           455           460
153 Pro Gln Pro Gly Thr Val Val Arg Met Gln Gly Leu Ala Tyr Asn Thr
154 465           470           475           480
155 Gly Val Lys Glu Ile Leu Asn Phe Phe Gln Gly Tyr Gln Tyr Ala Thr

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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000
 TIME: 10:45:55

Input Set : A:\Seqlist.txt
 Output Set: N:\CRF3\11082000\I697206.raw

```

156                               485           490           495
157 Glu Asp Gly Leu Ile His Thr Asn Asp Gln Ala Arg Thr Leu Pro Lys
158                               500           505           510
159 Glu Trp Val Cys Ile
160                               515
162 <210> SEQ ID NO: 3
163 <211> LENGTH: 14
164 <212> TYPE: DNA
165 <213> ORGANISM: Synthetic
167 <400> SEQUENCE: 3
168 ttttgatcaa gctt
170 <210> SEQ ID NO: 4
171 <211> LENGTH: 42
172 <212> TYPE: DNA
173 <213> ORGANISM: Synthetic
175 <400> SEQUENCE: 4
176 ctaatacgac tcactatagg gctcgagcgg cgcgccgggc ag
178 <210> SEQ ID NO: 5
179 <211> LENGTH: 12
180 <212> TYPE: DNA
181 <213> ORGANISM: Synthetic
183 <400> SEQUENCE: 5
184 ggcccgctct ag
186 <210> SEQ ID NO: 6
187 <211> LENGTH: 40
188 <212> TYPE: DNA
189 <213> ORGANISM: Synthetic
191 <400> SEQUENCE: 6
192 gtaatacgac tcactatagg gcagcgtggt cgcggccgag
194 <210> SEQ ID NO: 7
195 <211> LENGTH: 10
196 <212> TYPE: DNA
197 <213> ORGANISM: Synthetic
199 <400> SEQUENCE: 7
200 cggtcctag
202 <210> SEQ ID NO: 8
203 <211> LENGTH: 22
204 <212> TYPE: DNA
205 <213> ORGANISM: Synthetic
207 <400> SEQUENCE: 8
208 ctaatacgac tcactatagg gc
210 <210> SEQ ID NO: 9
211 <211> LENGTH: 22
212 <212> TYPE: DNA
213 <213> ORGANISM: Synthetic
215 <400> SEQUENCE: 9
216 tcgagcggcc gcccgggcag ga
218 <210> SEQ ID NO: 10
219 <211> LENGTH: 20

```

Per new sequence rules, the only valid responses are: Unknown, Artificial Sequence, or scientific name (Genus/species)

See circled portion of Item 12 on Encl summary sheet

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000
 TIME: 10:45:55

Input Set : A:\Seqlist.txt
 Output Set: N:\CRF3\11082000\I697206.raw

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220 <212> TYPE: DNA
221 <213> ORGANISM: Synthetic
223 <400> SEQUENCE: 10
224 agcgtggtcg cgccgagga 20
226 <210> SEQ ID NO: 11
227 <211> LENGTH: 25
228 <212> TYPE: DNA
229 <213> ORGANISM: Homo Sapiens
231 <400> SEQUENCE: 11
232 atatcgccgc gctcgctgc gacaa 25
234 <210> SEQ ID NO: 12
235 <211> LENGTH: 26
236 <212> TYPE: DNA
237 <213> ORGANISM: Homo Sapiens
239 <400> SEQUENCE: 12
240 agccacacgc agctcattgt agaagg 26
242 <210> SEQ ID NO: 13
243 <211> LENGTH: 24
244 <212> TYPE: DNA
245 <213> ORGANISM: Synthetic
247 <400> SEQUENCE: 13
248 tcttgaaacc tccagacaca agaa 24
250 <210> SEQ ID NO: 14
251 <211> LENGTH: 24
252 <212> TYPE: DNA
253 <213> ORGANISM: Synthetic
255 <400> SEQUENCE: 14
256 ggagatggta gacactggtg gagt 24
258 <210> SEQ ID NO: 15
259 <211> LENGTH: 24
260 <212> TYPE: DNA
261 <213> ORGANISM: Synthetic
263 <400> SEQUENCE: 15
264 tcttgaaacc tccagacaca agaa 24
266 <210> SEQ ID NO: 16
267 <211> LENGTH: 23
268 <212> TYPE: DNA
269 <213> ORGANISM: Synthetic
271 <400> SEQUENCE: 16
272 aagttacgat ttgcttcac tgg 23
274 <210> SEQ ID NO: 17
275 <211> LENGTH: 9
276 <212> TYPE: PRT
277 <213> ORGANISM: Synthetic
279 <400> SEQUENCE: 17
280 Phe Leu Gly Glu Phe Ala Thr Asp Ile
281 1 5
283 <210> SEQ ID NO: 18
284 <211> LENGTH: 9

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Please correct this error in subsequent sequences

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/697,206

DATE: 11/08/2000

TIME: 10:45:56

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\11082000\I697206.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date